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Sequence Listing was accepted.

See attached Validation Report.

If you need help call the Patent Electronic Business Center at (866) 217-9197 (toll free).

Reviewer: Anne Corrigan

Timestamp: Fri Sep 14 16:03:55 EDT 2007

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Application No: 10590633 Version No: 1.0

Input Set:

Output Set:

Started: 2007-09-04 12:29:12.260
Finished: 2007-09-04 12:29:12.868
Elapsed: 0 hr(s) 0 min(s) 0 sec(s) 608 ms
Total Warnings: 5
Total Errors: 2
No. of SeqIDs Defined: 6
Actual SeqID Count: 6

Error code	Error Description
W 213	Artificial or Unknown found in <213> in SEQ ID (1)
W 213	Artificial or Unknown found in <213> in SEQ ID (3)
W 213	Artificial or Unknown found in <213> in SEQ ID (4)
E 257	Invalid sequence data feature in <221> in SEQ ID (4)
E 257	Invalid sequence data feature in <221> in SEQ ID (4)
W 213	Artificial or Unknown found in <213> in SEQ ID (5)
W 213	Artificial or Unknown found in <213> in SEQ ID (6)

SEQUENCE LISTING

<110> ZACHARIAS, DAVID ALAN

<120> FUNCTIONAL GENOMICS AND GENE TRAPPING IN HAPLOID OR
HYPODIPLOID CELLS

<130> UFRF1100-1 (055932-0203)

<140> 10590633

<141> 2007-09-04

<150> PCT/US2005/06309

<151> 2005-02-25

<150> 60/548,509

<151> 2004-02-26

<160> 6

<170> PatentIn Ver. 3.3

<210> 1

<211> 18

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
Peptide

<400> 1

Met Leu Cys Cys Met Arg Arg Thr Lys Gln Val Glu Lys Asn Asp Asp
1 5 10 15

Gln Lys

<210> 2

<211> 17

<212> DNA

<213> Mus musculus

<400> 2

gtccccaggtc ccgaaaa

17

<210> 3

<211> 68

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Consensus
splice donor site

<400> 3
ccgctcgaga cttacctgac tggccgtcgt tttaagacga gctccctagc tagtcaggca 60
ccgggctt 68

<210> 4
<211> 6
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Consensus
motif

<220>
<221> MOD_RES
<222> (3)..(5)
<223> Variable amino acid

<220>
<221> MOD_RES
<222> (6)
<223> Ser or Thr

<400> 4
Met Gly Xaa Xaa Xaa Xaa
1 5

<210> 5
<211> 4
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Illustrative
CRD motif

<400> 5
Asp His His Cys
1

<210> 6
<211> 13
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: C-term tail
peptide of GFP: Yck2p

<400> 6
Lys Ser Ser Lys Gly Phe Phe Ser Lys Leu Gly Cys Cys
1 5 10

